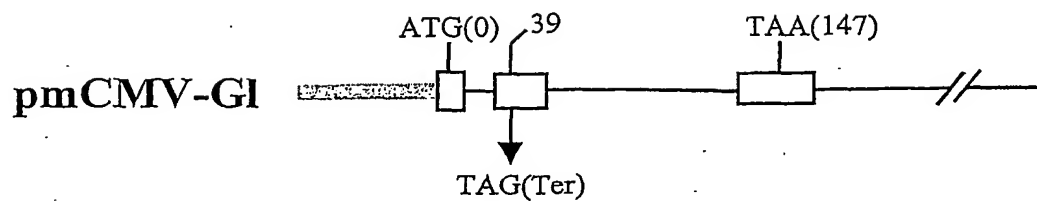


A



B

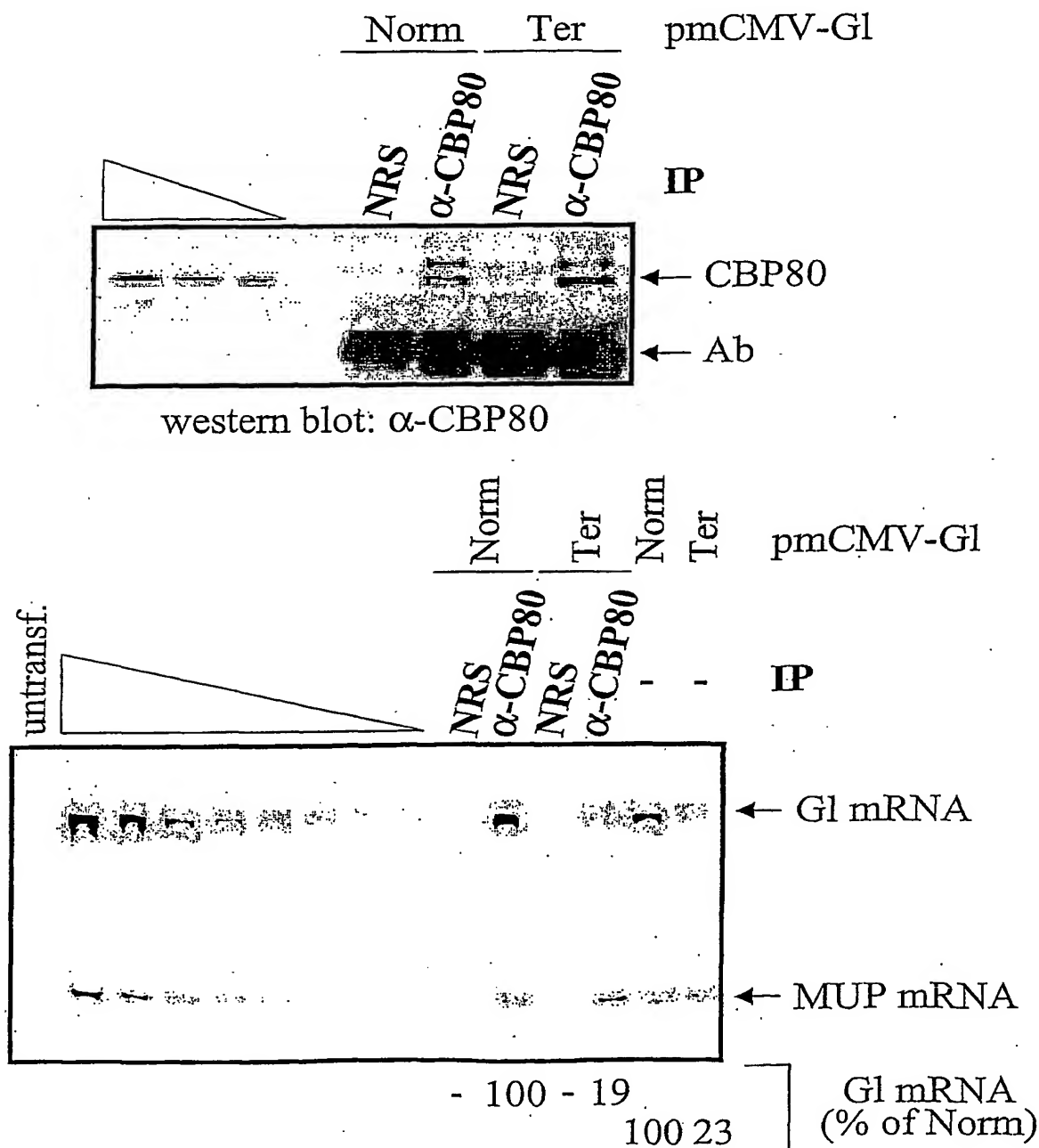


FIG. 1 BEST AVAILABLE COPY

		Norm		Ter		Norm		Ter		pmCMV-Gl			
		α -CBP80		α -CBP80		α -eIF4E		α -eIF4E		IP		% of cell protein in IP	
		NRS	-	NRS	-	NRS	-	NRS	-			α -CBP80	α -eIF4E
8x10 ⁵ cells										α -CBP80		34±2	0
										α -eIF4E		0	15±1
										α -CBP20		26±2	0
										α -eIF4G		2±1	10±4
32x10 ⁵ cells										α -PABP2		11±1	0
										α -Upf 1		0	0
										α -Upf 2		9±3	0
										α -Upf 3		7±1	0
										α -rpL10		2±1	14±3

		Norm		Ter		Norm		Ter		Norm		Ter		pmCMV-Gl	
		α -CBP80		α -CBP80		α -eIF4E		α -eIF4E				IP			
		NRS	-	NRS	-	NRS	-	NRS	-						
		-100		-17		-100		-12		100		20		Gl mRNA (% of Norm)	

FIG. 1

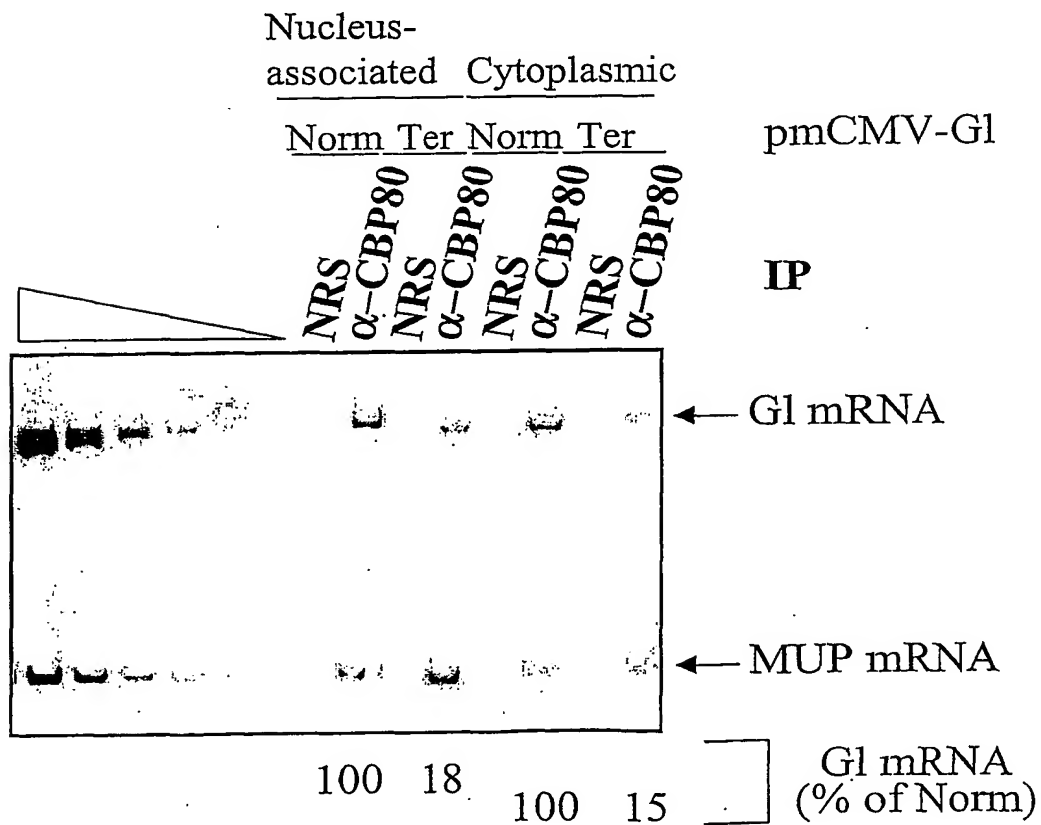
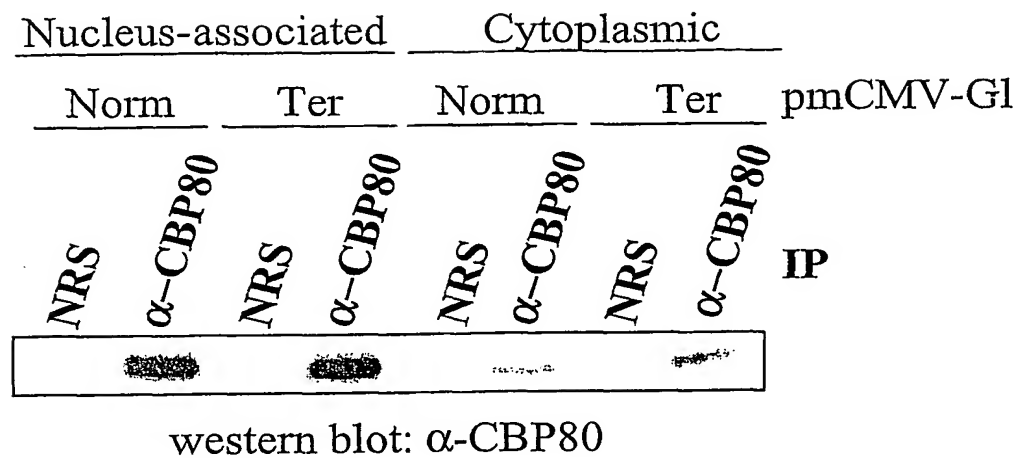


FIG. 2

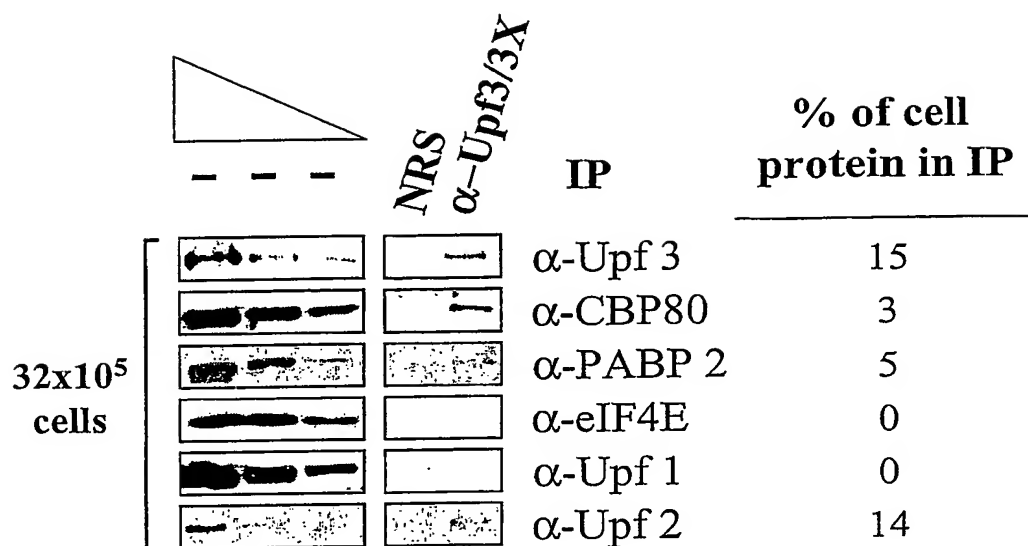
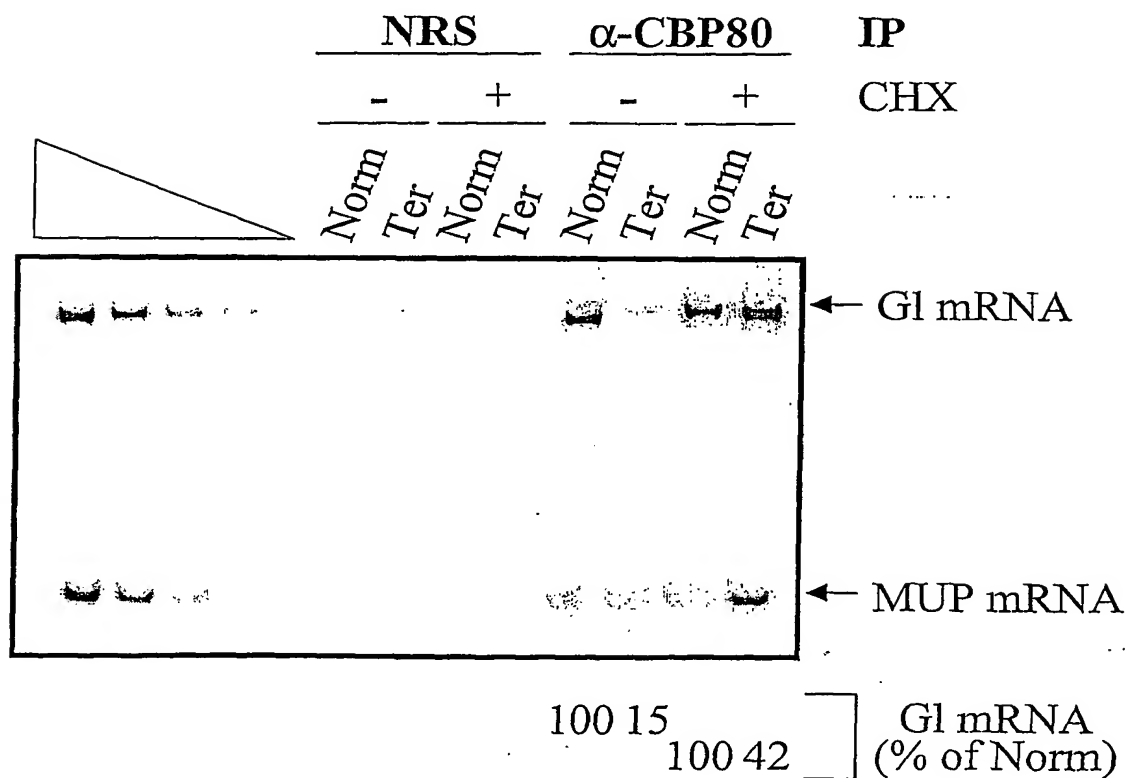
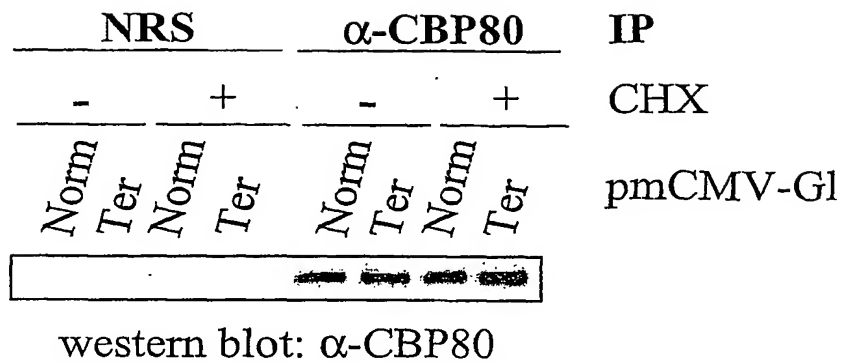


FIG. 3

A**FIG. 4**

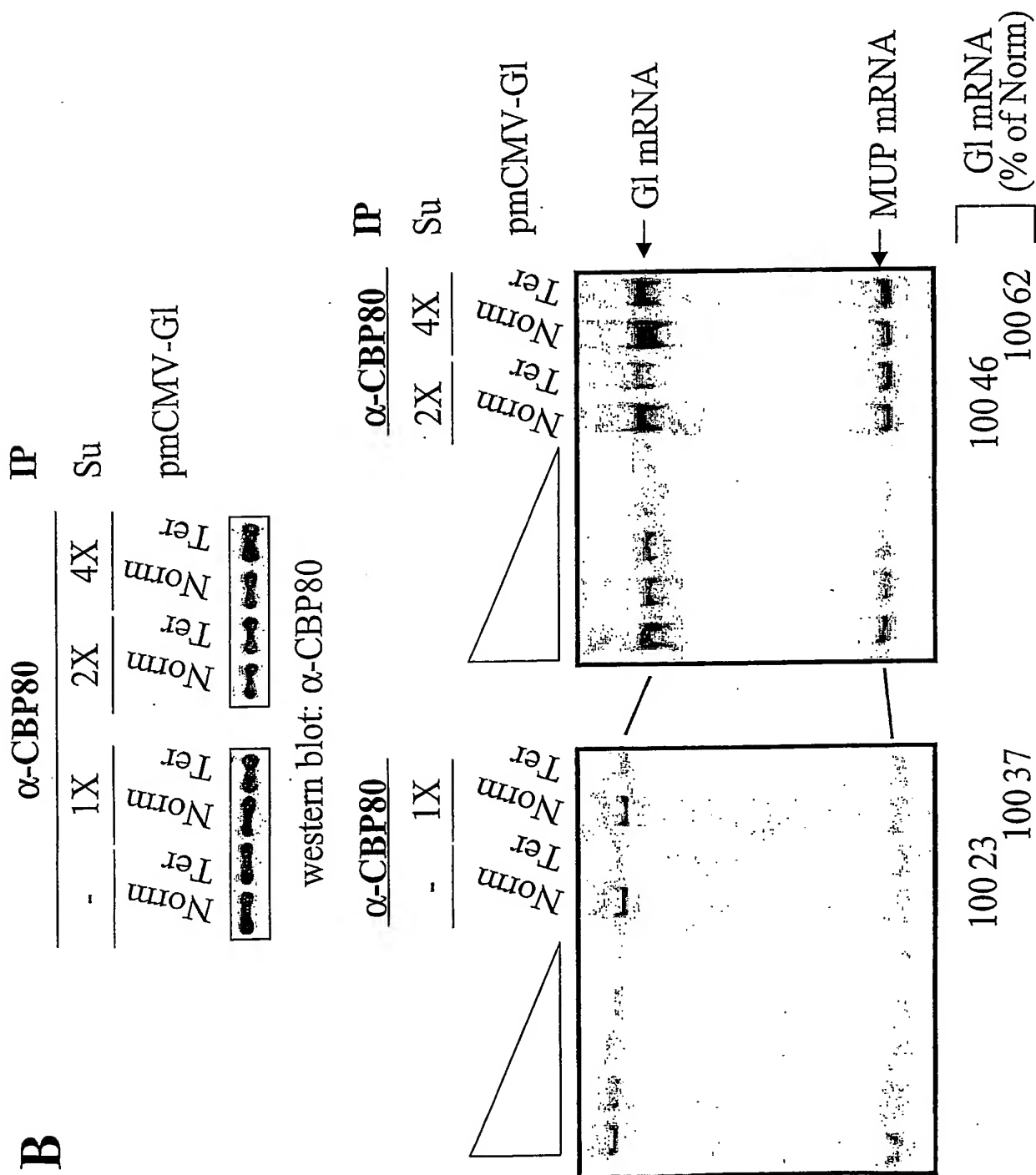


FIG. 4

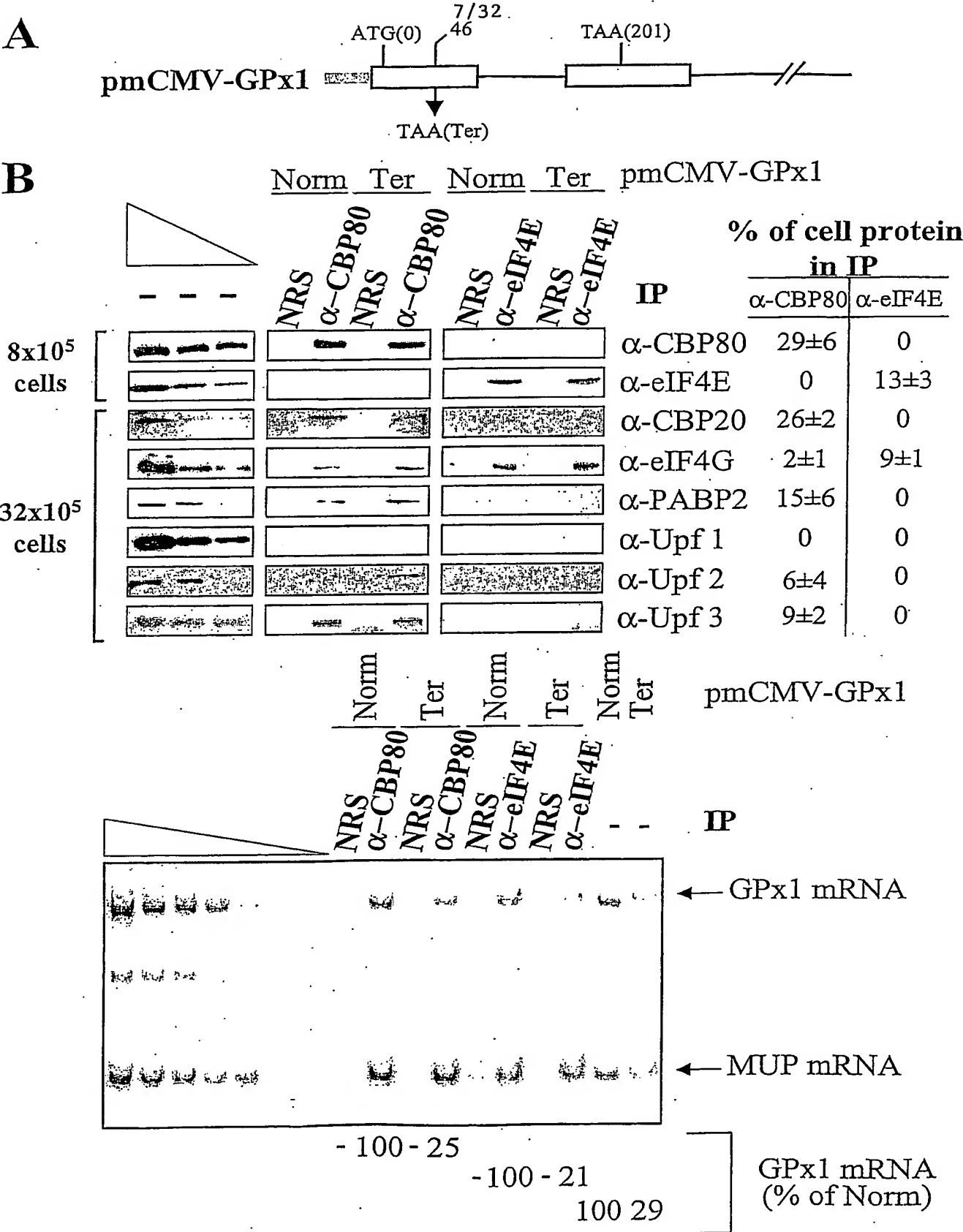
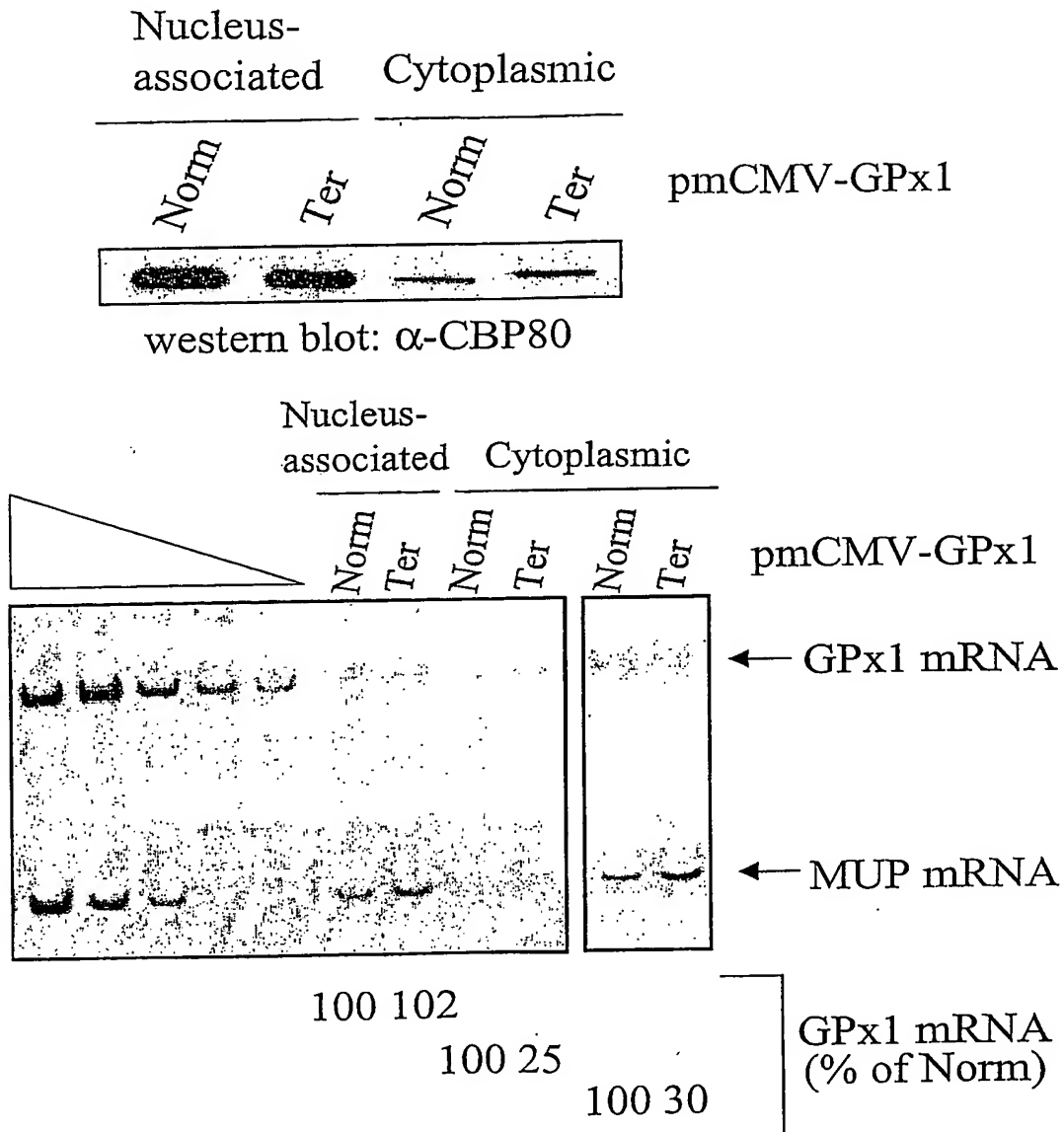


FIG. 5

**FIG. 6**

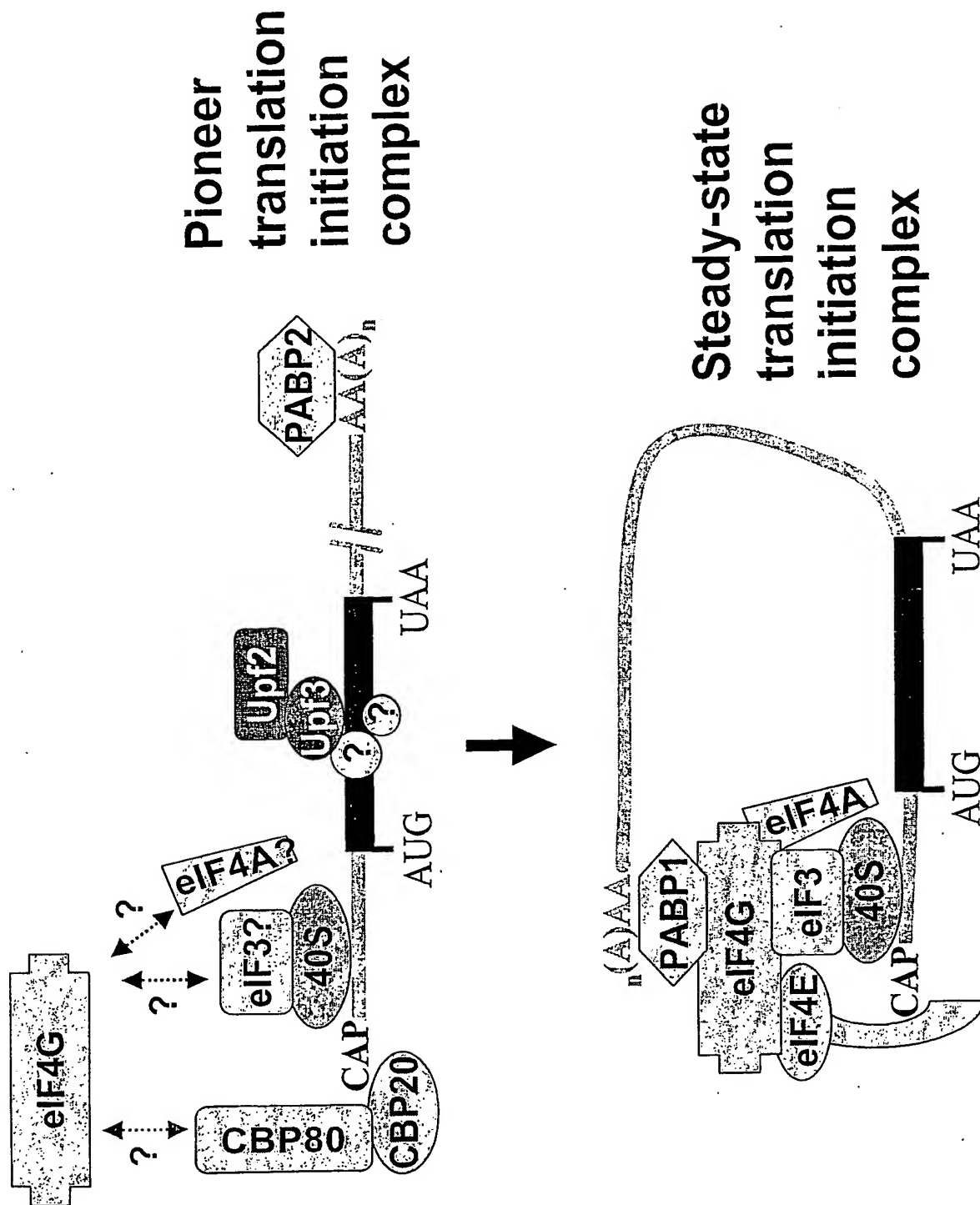


FIG. 7

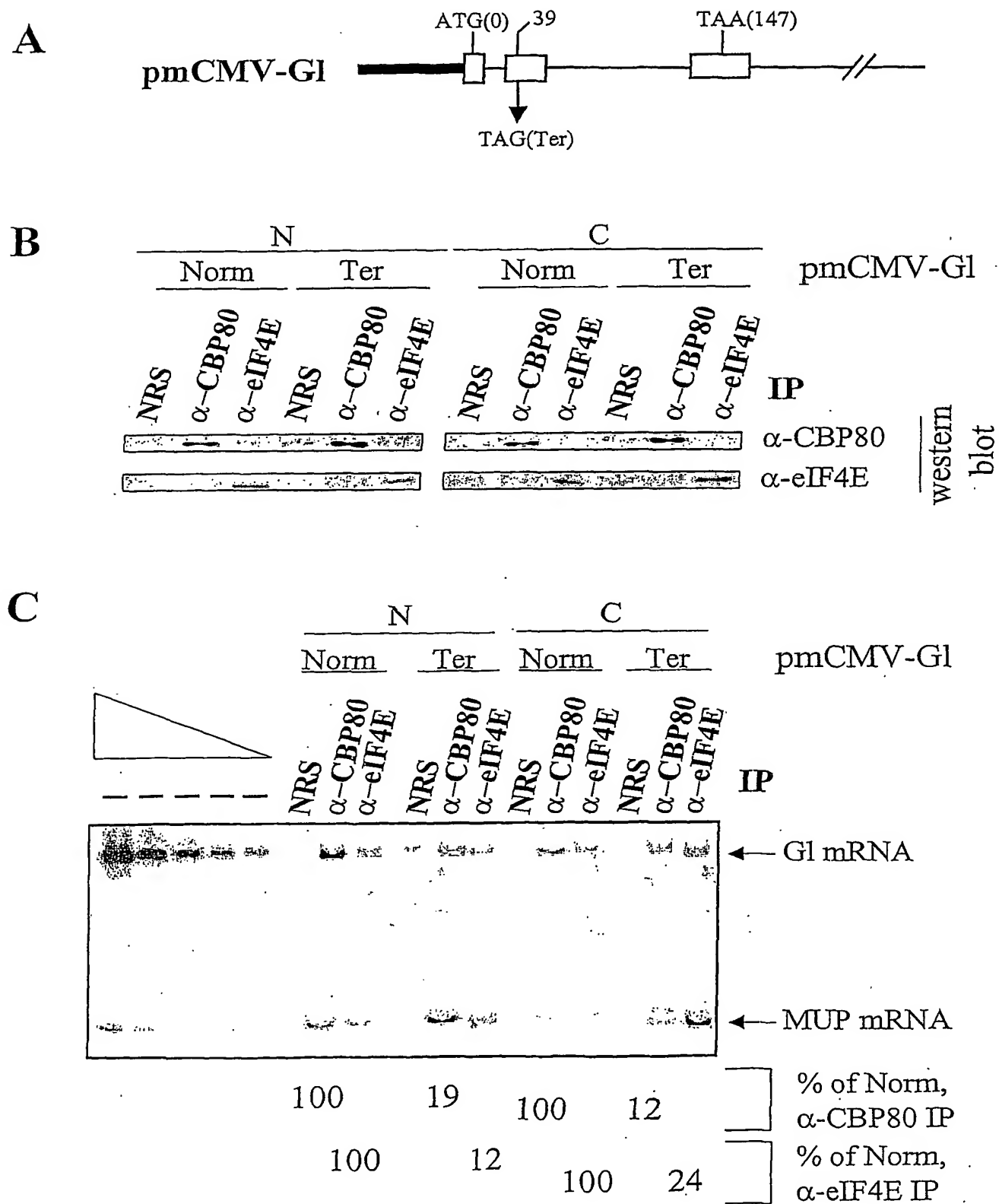


FIG. 8

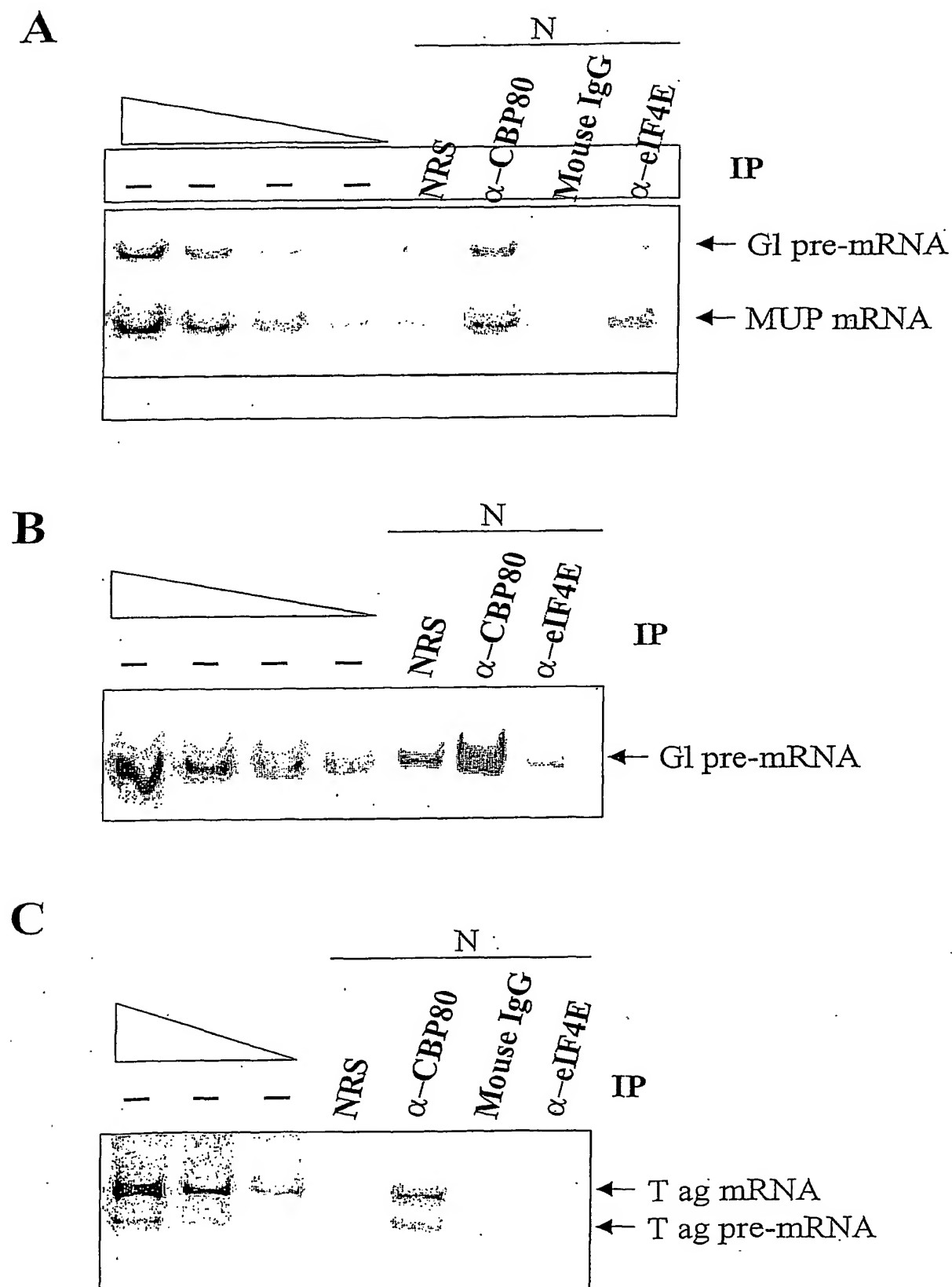
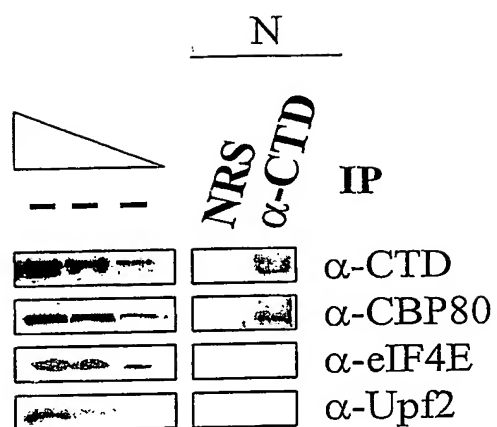


FIG. 9

**FIG. 10**

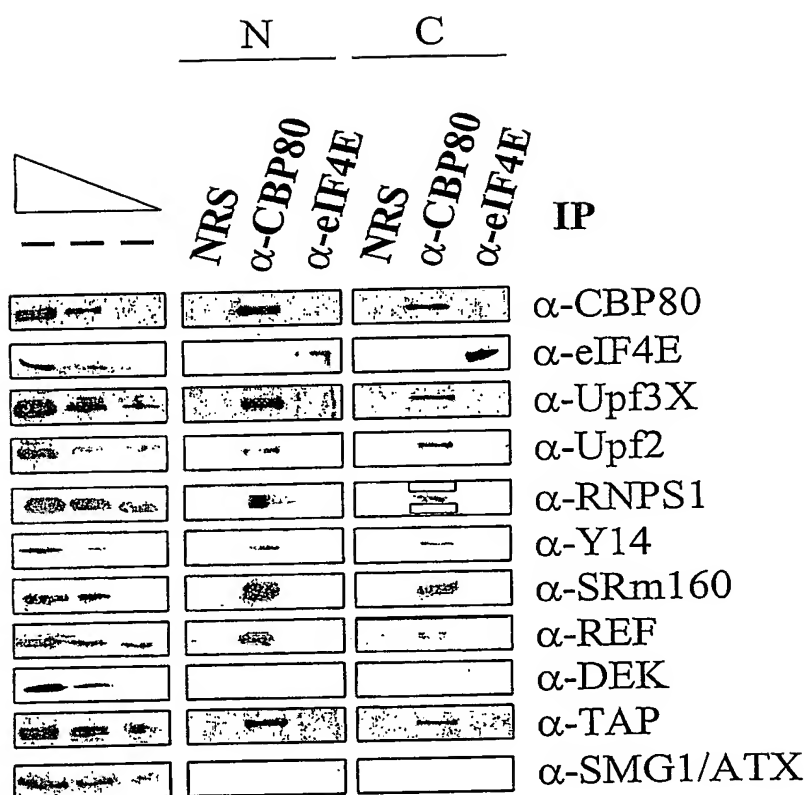


FIG. 11

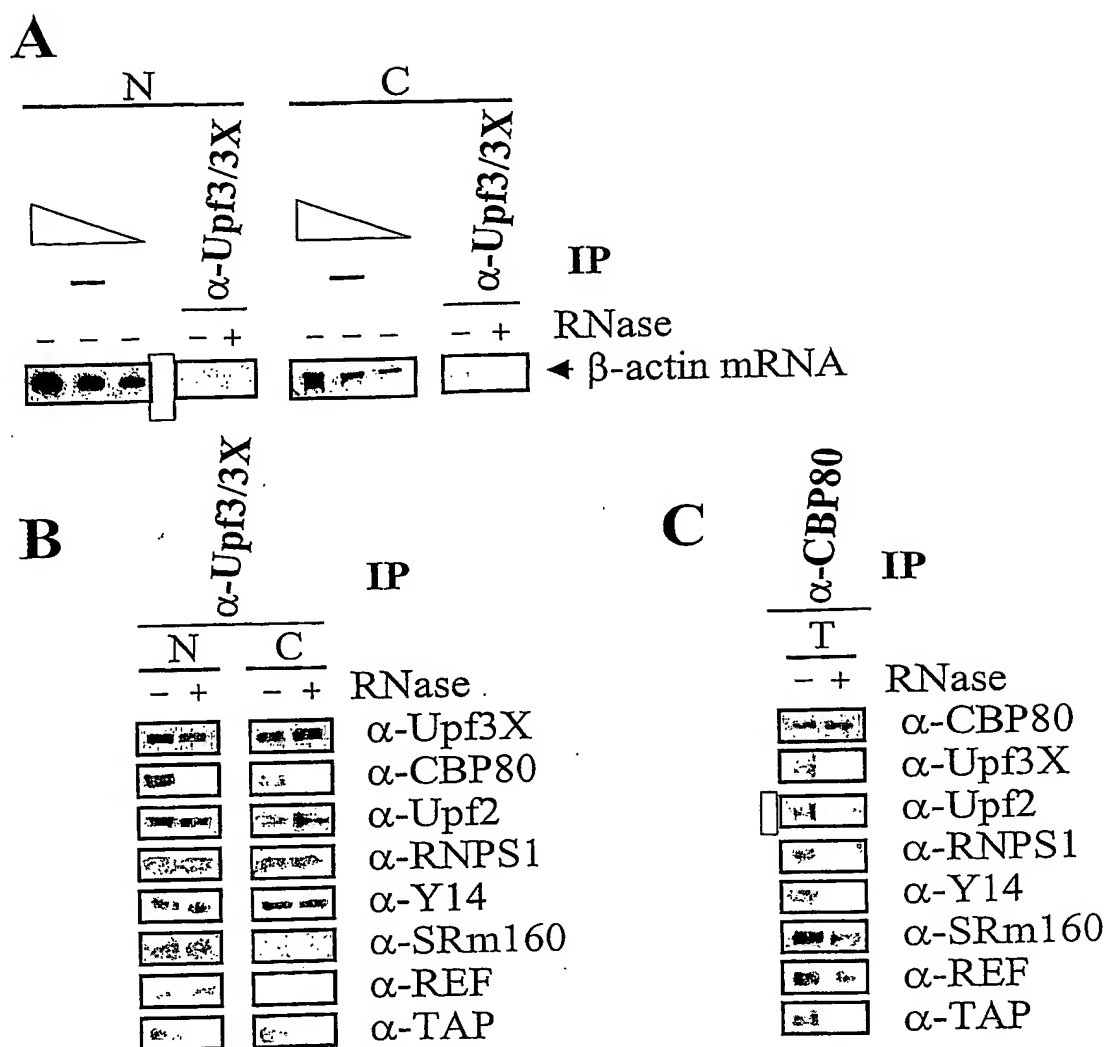


FIG. 12

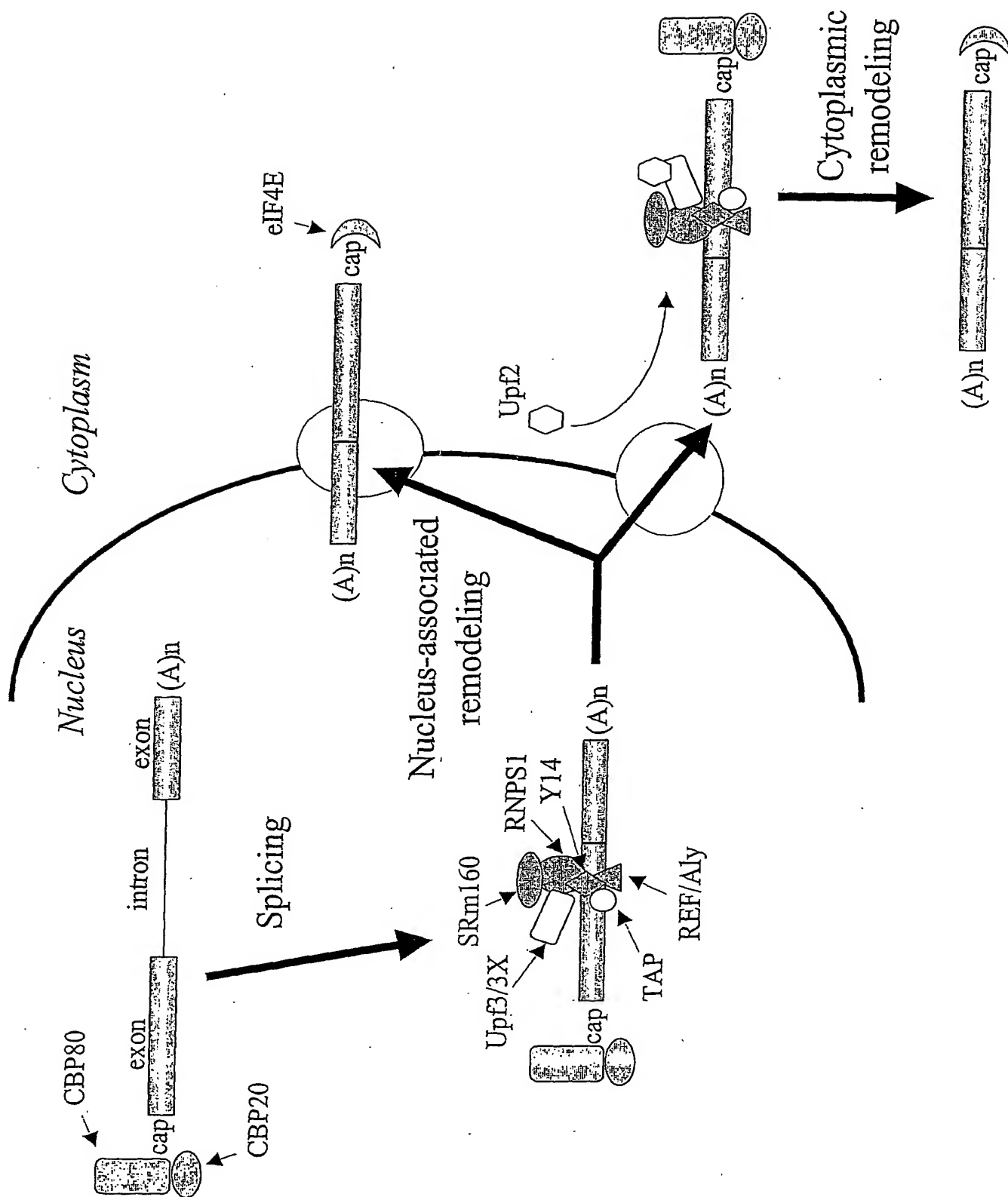


FIG. 13

MAEGLERVRI SASELRGILATLAP QAGSRENKMKELKEP <u>FRPK</u> ONRRPDLEIYKPGLSRLRNKPKIKEPPGSGSEEPKDEIVN :	80
DRDCSAEVENGVQVQVQVQCKELNNOEQNGPFDIPENNRRGQSEFPRTAQDBRSLAIKIKTKKPPDQJLYQVQGRRLQVTSKESA :	160
SRVEEVEVLNQVQQLRVBEDECRCNVAKAEVANKPDRAEIKSPGGGRVGLAKIEKKGKRMKGQGEVTRHTDDPARGPPGS :	240
AKRYSRSDKRRNNRYRTRSTSGAGNSAEGAGLTLDGQ <u>RRRR</u> DDTTERKPPKKQVSVSSTDSLEDRIDEPFDGLPPRS :	320
SE <u>RRKR</u> RLERNWSGRCGEQEKTSKAEYRGTRTLVFDAAEMKNKSPMVSRAVDMDRGPEKGLSSGGKSGEKQSKNPKQE :	400
LRGRGRGLILPAHLTILSVNSAGSPESAPLGPRLLLFGSGSGKCSRHWGRGGTTRRLWDPNNPDPQKPAIKTKTQQLHFLDIT :	480
DEVSTPSWGDSRQAQASYKQFNSNDYYYPRTPGPASPYPYTGYNLQYVPVTNGVYGPYGPYPTPGGQYVCSPLP :	560
TSTMSPEEVEQHMRNLQQLLHRLVRADNQELQNLNLSRDRISPEGLEKMAQRAEQLLQYRCILLDITFSDNNQVND :	640
QILWKNAYQYQVIEKFRQLQVLDPVENPEQIRNRLLELDEGSDFFDSLLQQLLVYKFKLEDDMDGLAIRSKPLRKTVKY :	720
ALISAQRCMICOQDTRARYEQASDTPYVYKGRSWYLKQAHIAKGRPNYNOQLLAVYTRKLDADVYTYMRSLSANPNLI :	800
TAKESLMSLFEETKRAEQAQMEKKQHEEFDLSPDQWRKGGKSTFRHVGD DTTRLLEIWIHPSHPRSFMFSGKDSQENGL :	880
GSLSPDLNKRAPFLISFLAHAGKLFTRIOMETPFAVAKVLEKEFPQVLLHQSPSIPGSTRMLQMTNMFAVHNSQLKDCFS :	960
EBCRSYIQEQAALLGAMFLSLVRRCTKLKESAKAQLSSPEDQDQDDIKVSGFPDTEKLLPSVKVWSDMWLGYPDWT :	1040
NPPPTSLDPSHVAVDVWSTLADF CNILTAQVQSEVPLYKOPDDMLLLILEEDRLSSGFPFLAABPDPCYVEKTSKIV :	1120
IADCKRVTVLKYFLAELCQGERPILLAFKGGKYVSAVPVPTDGMKEMSGEGTLEDEEEDVIEDFEEDSAEGSGGBD :	1200
DIRELRAKILAKARKIAEQQRQREKIQAVLEDHDSQMRQMELEIRPL <u>FLVPD</u> TNGFI <u>DHLASLARILLES</u> SRKYILVVP <u>LI</u> VI :	1280
NELDQLAGQOETDHRAGGYARVVOEQAKRSFTEFLQRPBSRSDCIRALTRSGNELESIAFSDSETITGQLGNNDLILSCC :	1360
LHYCKDAKMPKASKEPTIRLLREVALLTDDRLNRVLAIRTNVPVNDIPAFLTWAQVG :	1419

CCTGGCTCGCGCGCGCGCGCGCGGAGACCTACAGCTCGATAGCAGCAGCCGAGGAAATGCGGAGAGGCGCTGCAGCGCTGTGCGGATCTCCGCGCTCGGAGCGTC	1
CGCGGATCTCTGGCTACTCTGGCGCCCGCAGCGCGCGGAGCAGGCAAAATATGAAAGGATTAAGGAGGCGCAGCGCCGCAAGATACAGCGCGCTCCAGATC	200
TGGAAATCTTAATAGGCTGCGCTTTCTCGGCTAAGGAAACAGGCCAAATACAGGAAACCCCTCGGAGAGTGAGGAGANTCAAAGATAGAAATTTTAATGACCG	400
AGATTGCTCTGTCGTTTGAAATATGCTACACAGCCGCTTAAGATGCTCTGAAAGGAACTGAAACAACAGCAGAGCAGAGATCGCTCTAAGACCCAGGAATAATAT	600
CGGGACAGAGAATCTGCTCTTCCAGGACTGCTGCGACAGGAGATCTGAGTCTAARAATTTATCAAAGAACCAAGCACTCGAGATCTATCAGCTGCT	800
GCAGCTGTTCTGAGATCTGTAGAAAGATATCCGCAAGCTCGGTCGAGGAGAGACAGATCTCAACCAAGTAGAAACACTGAGATGATAGGAGAAATGATGT	1000
TAGGGAAAATTTGCTCGAAAGGAGGTTGCGAAATAAACCCAGCAGCGCGAGATAGAAAGAGCCAGGTCGTGGGAGATAGGGGCTGCTCAAAGAGAGAA	1200
AAAGGAGAAAGAGATAGTGGAAAGGGGGGGGGTAGGAAACCCACAGCAGCCCGCGCGCGGAGGCGGCGCTCGCAAGACGATCTCCGCGCTCAGACA	1400
ACAAGAAAGATCGCTACGCGCGCGGACGACAGCTCAGCTCAGCTGAGTGCAGCAACAACAGCTCTGAGGAGAGCTGCGCTGACGAGTATGAGTTCTCGCGCCGCG	1600
ACAGGATAGAGACCAAGAGAGAGGCCACCACTGGAAGAGCAAGTGTCTGTCTCAACGATCTCTTGAAGAGAGAGCAAGATTGATGAGCTCTGATGATAT	1800
GGACCCAGGAGAAATTCAGAAAGAGAGAGACTTTAGAAAGAAATCTGCTCTCGCGCTGGGAGGGTAGCAGAGAAACCAAGTCTAAAGAAATCTAGGCGCA	2000
CTCTCTCGTGTGATCTTGATGTCAGAGACCATTAACAAGAGATCTCCCACTGGTAGGTCAGCGAGGATGATATGGATAGGAGAGAAAGCTCAGAAAGAGAT	2200
GACAGTGGGGGCGAAAGCTCTGAGAGAGCAGGATGCGAAATAACCGAAACAAGAAATCTCGGGTCTGGTGTGTGGTATCTGATTTTGGCTCGCCGATATCC	2400
ACCTTACTCTCAATTACAGCGAGTCTCCAGAGTCTCGCGGCTTTGGGACCTCGGCTTTGTTTGGATCTGATGATTAAGGATCTCCGAGTGGGGGCGCT	2600
GAGGACACACACCGCGATTTGGGAGCCPAAACATCTGATCAGAAACCTGCTAAAGACTCAGAGCCCGAGCTACATTTCTGGACACTGATAGTGA	2800
AOTCAGGCTCATCTCTGGGAGTACTCAGCCGACGCTCAGGATCTACTATAGTTCTTAAATCTCTGACACCCCTATATTTATCCCGGACACACCGAG	3000
CGCTCTCCCAAGTATCCCTATACGGGCTATAAACCCTCTACATGACCAAGCTGGGAGCTGACAGATGGTGTGTATCCAGGCGGCTTACTACCGAGCTACCCGA	3200
CTCGGTACAGACAGATATGTGTGTATCGCTTACTCTACTCCAGCACATAGATCCCGAGAGGTAGTACAGACACATGAGGAACCTCTCAGCACACAGGATGACTC	3400
CAGGCTCTCCGGGTGGTGACACACAGAGATCTCAGCTCAGCAACTGCTCTCTCGAGGAGCGGATCAGTCCGGAGGCTCGGAGAGAGATGGGCGAACTC	3600
ADAGCTGAACCTGCGAGTATATGAGTATGCTGTATTTCTATTAGATATTGAGTCTCTGTATATACAGATATGGTGAATCAGATCTGATGCTCTGGAGAGATCTCTT	3800
ATCAGGTGATTTGAGAGATTCAGGCAACTGTCTAGAGTCTCGAATTTGAGAACCCAGACAGATGCGGAACACTTTTGGAGCTCTTGGATGGAGTAG	4000
TGACTCTTTTATATAGTTTGTCTCAGAGAGCTCAGGTTACTTACAGTCTCAATCTGGAAGACACATGATGTGTTCTGCCATCTCGCACAGCCATTACCG	4200
AAACGCTAAATATGCTGCTTGATCAGTCCGACAGGATGATATGCGAACGGAGATTTGCTAGATACGGGAGCAGACCGGATGATACAGGAAATATTG	4400
GGAAAGAGCCAGCTTGATACAGTGAAGCCAGCACATCTGCTCCAGAAATGCGGCGCCCTATCAACAGTTGGCTTTCTGTCAGTGTATATCAAGGAGGAA	4600
CTTGAGCGCTGTTACTACTATATGGTCAGTTTASCTGAGCAACCTACTCTGACTGCCAGAGAGACTCTATGAGCTCTTTGAAGAGACACAGCG	4800
AACCTCAGAAACAGATGGAAGAGAGCAATCAGGAAATTTGACCTCAGGCGCTCAGCAGTGGCGGAAGAGAAAGTCTACTTCTCCGAGTGTGGAGATG	5000
ACACCACTGCGCTGAGGATCTGGATCTGATCATCTCATCTCAGGCTCTTCCAGAGCTCAGTGTAGGAGAGATCTGATCAGAGAAATGGGCTGGGACG	5200
CGTGGATCCGATGATTTGAACAAAGGTTTACTCTCAGTTTCTCTCATGCCCCATGGGAAGCTGTATACCGGATTTGATAGAGACATTCCTATCAGT	5400
CTGAGAAAGTCTCCAGAGGATTCAGGTTCCAGTGTCTTACCTCAGCAGCTGCGCTCCCATTTCCAGATATCCCGCATCTGGGCTATGACAGTCAAGTTTGT	5600
CATGACACATCTCCGACGAAAGCTCTCTCTCGGAGGAGTGCGCGCTCTGTAATCCAGGAACAACGCGAGCTCTGGGCTTGGCCCATGTTTCTCTCAT	5800
GCTGCCGCTGACCTGCTTCTTAAAGATCTCGGCAAGCTCAGCTGCTCTCTCTCGAGGACAGGATGACAGAGACATCAGGTTCTCTCTCTT	6000
CTCCGAGACCTGAGGAGTCTGCTCCGAGTCTTAAATCTGTGTCAGATTGAGATGCTGCTCAGCCGATACCTGGATCTCTCCCTCCATACCTCCAGT	6200
TGCCCTCGCTGTCTGTAGTATGTGTGCGACCTGCTGCTGATTTCTGTATACCTGACTGCTGATGATCAGTGTAGGTTGCCATCTCAACAGAGACC	6400
GGATGATGATCTCACCTCTTACTCTGAGAAAGATGGGCTCTCTCGGCTTTGTCTCCCTCTGTGGCTGCCCATCAGAACCCCTCTACTCGTGAGAGT	6600
ACCTCGGATAAGCTTATTTCAGTCTGAGAAAGGTCACAGTCTGAGATTTCTTGTGAAGCCCTTTGTGGAACGAGTCTTCTGCTGATCTTCA	6800
AGGTTGGAAGATATGCTCTCAGCTGCCACCTCCGACACCATCGGAAAGAAATGGGAAACACAGAGGAAACAAGCTCAGGAATGAGGAGAGT	7000
GTGTATGACAGCTTTCAGGAAATTTACAGGCTTGAGGCGAGGCGAGGAGAGATCATACGAGGATCTCGGCGCAAGACTCGCTCTCGCCGAGAG	7200
ATAGCTGAGCAGCAGGCTGGCAGGAAAGATCAAGCTGCTCGAGGACACCATCAGATGAGGCGATGACATCTGAAATCAGACCTTTGTTCTCTG	7400
TACGAGACACCAAGGATCTTATGACCACTCAGCAGCTCTGCGCGCTGCTGAGAGACAGAAAGTACATCTCTGTGTCCTCTACTGATCTGATCTG	7600
GCTGACAGGCTCGGCGCAAGGGGACAGGAGACAGACACCTCGGCTCGGGGCTACCGGCTGTGTGATCAGAGAGAGGCGGCAAGTCTCATGAGTTCTCTCGAG	7800
CAGGAGATCTCGAAGTCCGAGCTCTTGCTCGGAGCCCTCAGCAGCCTGGGATGAACTGAACTGACATCCATGCTCTCCGACGAGGACTCATCTCGGACG	8000
TGGTAAACAGAGATCTCTTACTCTGCTCTGCTGCTGCATCTGACAGCAAGAGAGGCTAAGAGATCTTATCGGCGCAGGAAACAGAGGAGCAATCTCGGCT	8200
ACTGTCGGAGGGTGTGCTGTGTCAGGATGACCGGAAACATCTCGCTGTAGAGGCGCTCAGCAGAGATTTCTGTTATCGGGAATCTCCAGCTCTCTCATCGG	8400
CGCCAGGTGGGCTGGAGGAGCCACATGGGCGCCCCCGCTGGAACTCTCTGAAAGGCAACAGGCGGCCAGCTGATGACGGAAGATCTGACGCT	8600
CTGCTGAGCAGCAATCTACCCAGACATTAATAACTCTGCTCTTCCAAACGCGCCAGCAGGAGGAGTCTGTGGGGAGACTGCTCTCTCAGCAGGCGCTCCAG	8800
GATCGGAGAGAGAGCTGCTGGGAGCCCTCTGGGCTGTCAGAGTTAGCAGGAGATGGTGGCTGACAGCAACAGACTGGGACAGGACAGATAGGCGCCGC	9000
ATGCTCTCAGCCTCTCTCCCTCCCCCTCTCATCTCAAAGGCTGAGGAGGGGCTCTCGCTCGGGAGCGAGCACTCTCTCAAGTGGAGACAGGCGAG	9200
GCTTCAAGTTTTCGTGACAGTCACTGAATATCAAGTCTCTCTTCTCATCTCAGAAACCTCTGCTGTGAGATGTTCTTCAAGGCTTTTGGTTAAGTCAAGAG	9400
AACTGTCGCGAGGCTGTGTGCTCCGAGCTCTCTCGAGGCTCGACCTCGCCCATGAAACCAAGTCCGCTCTCAGACAGATGTCTCTTAAACAGCAGCTCTGGG	9600
CCGCACTCTTCTATCACTACACCCAGCACTTACACACACCTCTCTGCTGCTCTCTCTGCTCTTCAAACTGGGTCCTTAACTCTGCGAGTGT	9800
CGTCTCGGAGGACATGATCTAGATCTAGAGGAAAGAGGACATCTGCTCTGAGGCGGCTGCTCTCGGCTGGGCTCCACTCTCTCTCTCTCTCTCTCTCTG	10000
TTCTTTTCGCTTAGGCTCTCTCTCAGGAAATGTCTGACTCTCTCTCAGCTCCGCTTCACTCCGCTCTCTGCGGCGTACCTCCCTCTCAGAAATAGCCGCT	10200
ACCTCTCTCTCCCTCTCTCTGATGATCTCTTCTCACTCTCGATCTCCCTTCTTCTCTCAGCGGCTCTCTGCTTATAGGGGTTCTCACTCTGAACTTTCC	10400
CTCTACTACTCCATGCGAGGAACCTAGTACAGATCTTCCACAGGCGGCTTCCACCTGCTCTGCTCTGCGGAGAACTTTCAGGCTGACGAGCAGCCGA	10600
GCTCGAGGACTTCAGTCTGGGACCGCTCGCTTCAAGTGTCTCAGCTTCTGCTCTCTCCGCGGAAATCTGACATGACACAGGCGGCTTCTCT	10800
TGCTCTTTAGTGGTACTCTCAGTGGGAGGCTTCTTACCAAGATTAAGTCTCTGAAACCCAGGACAGACAGGACATCTTAGGGGAAAGACAGGGG	11000
TTCCGGTGGAGGCGGGCAAACTCTTAATGGGACAG	

FIG. 14

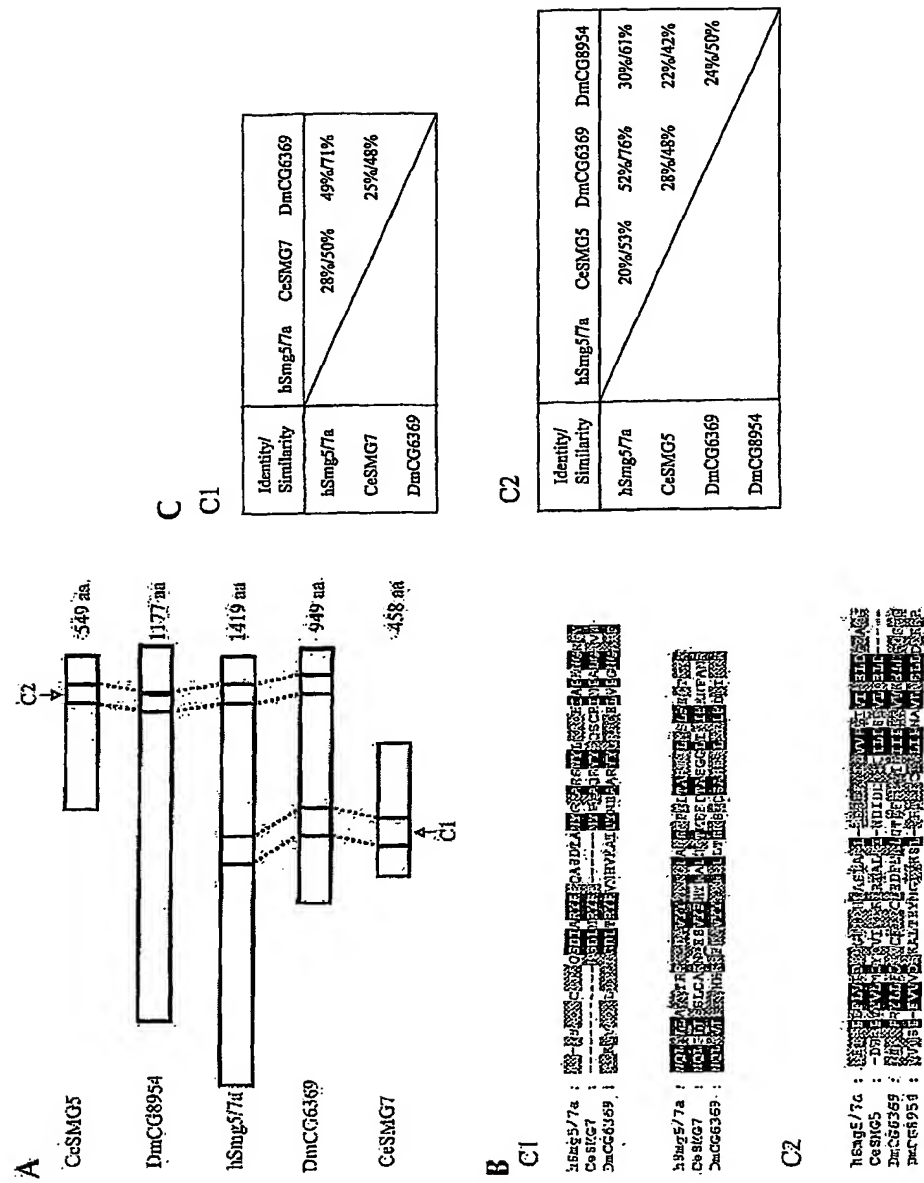


FIG. 15

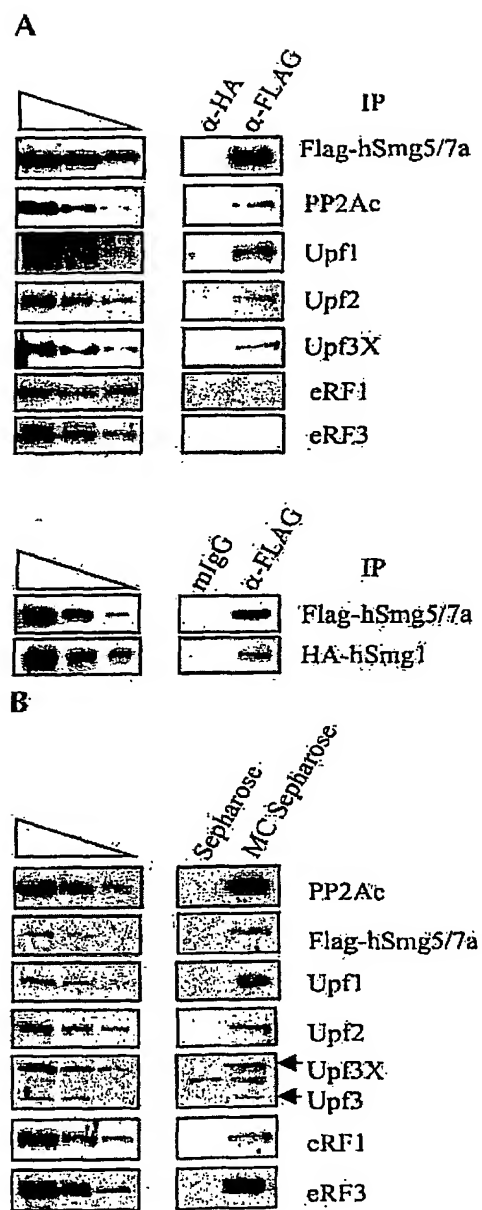


FIG. 16

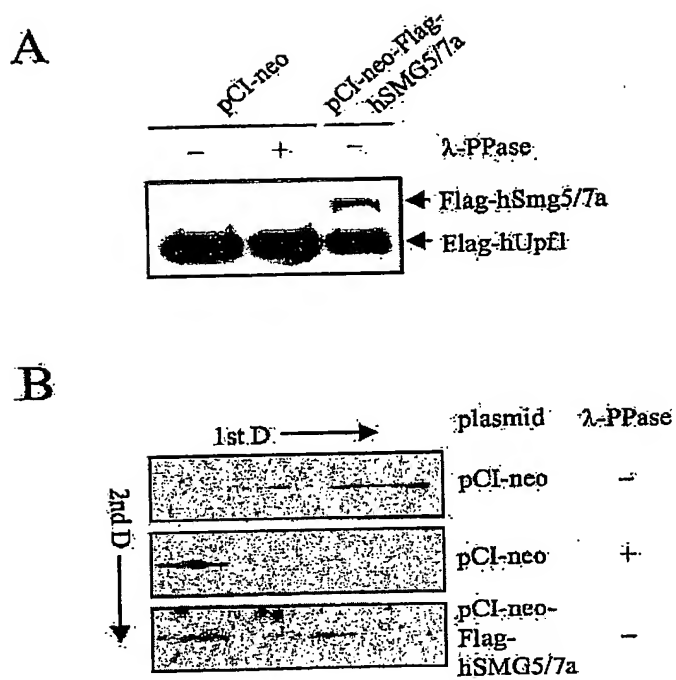


FIG. 17

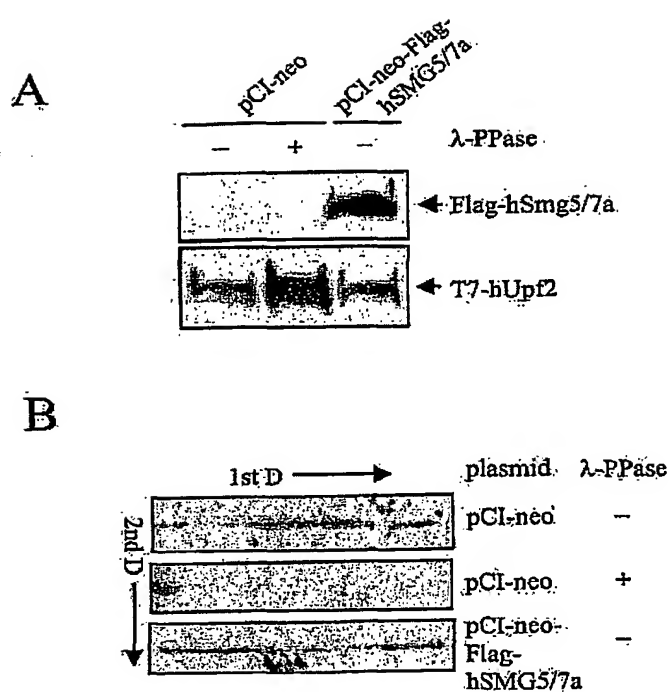


FIG. 18

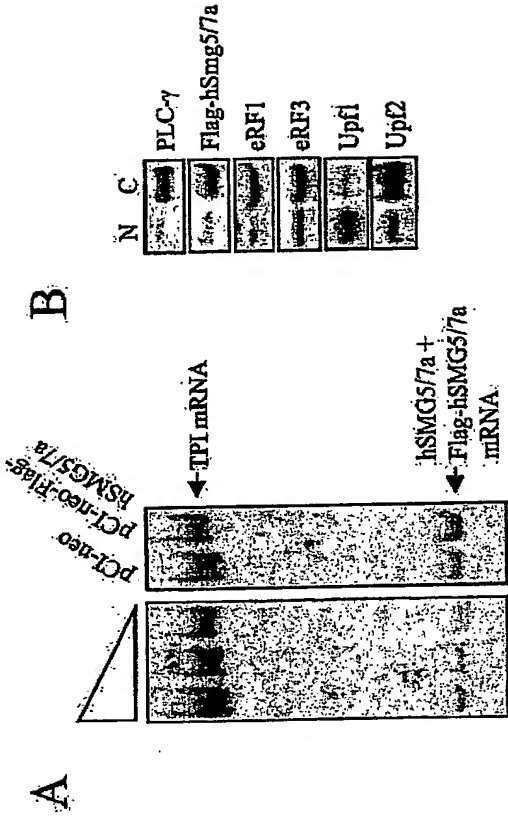
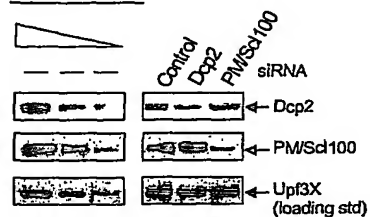
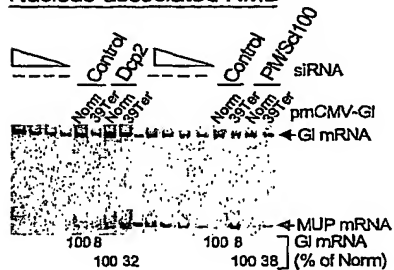
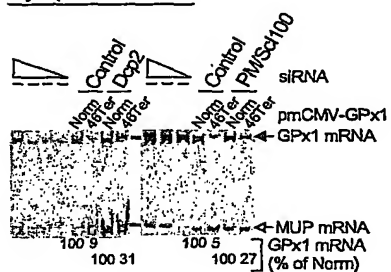
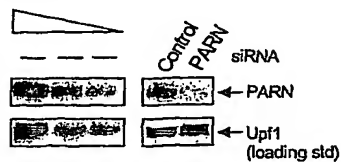
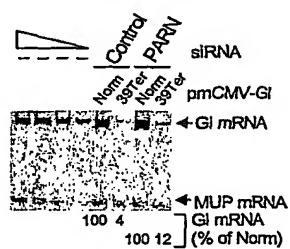
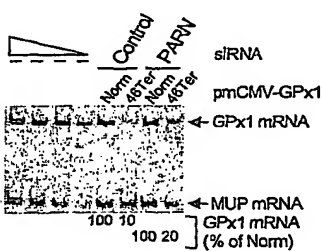
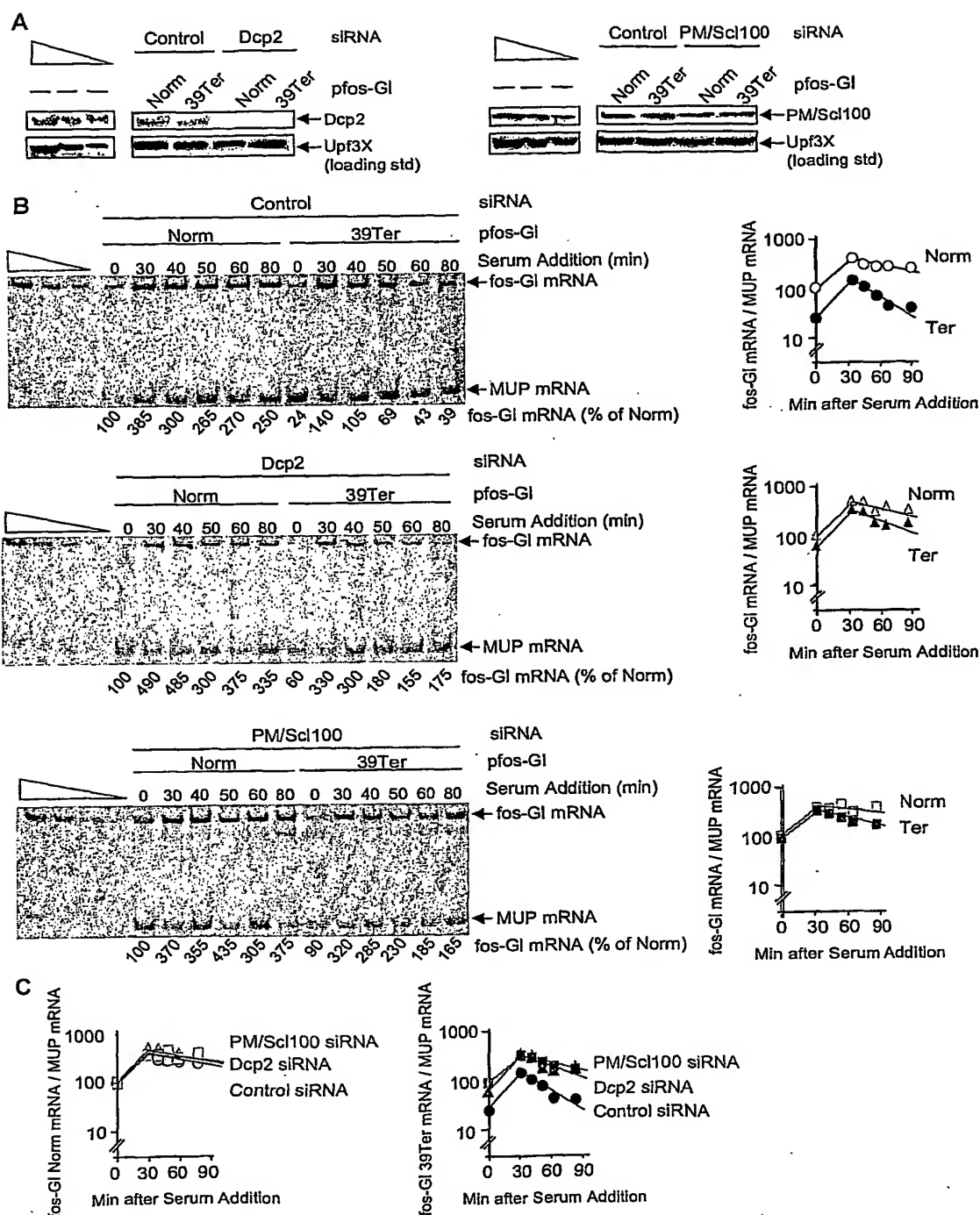


FIG. 19

A. Down-regulation of Dcp2 or PM/Scl100Western blotNucleus-associated NMDCytoplasmic NMD**B. Down-regulation of PARN**Western blotNucleus-associated NMDCytoplasmic NMD**FIG. 21**



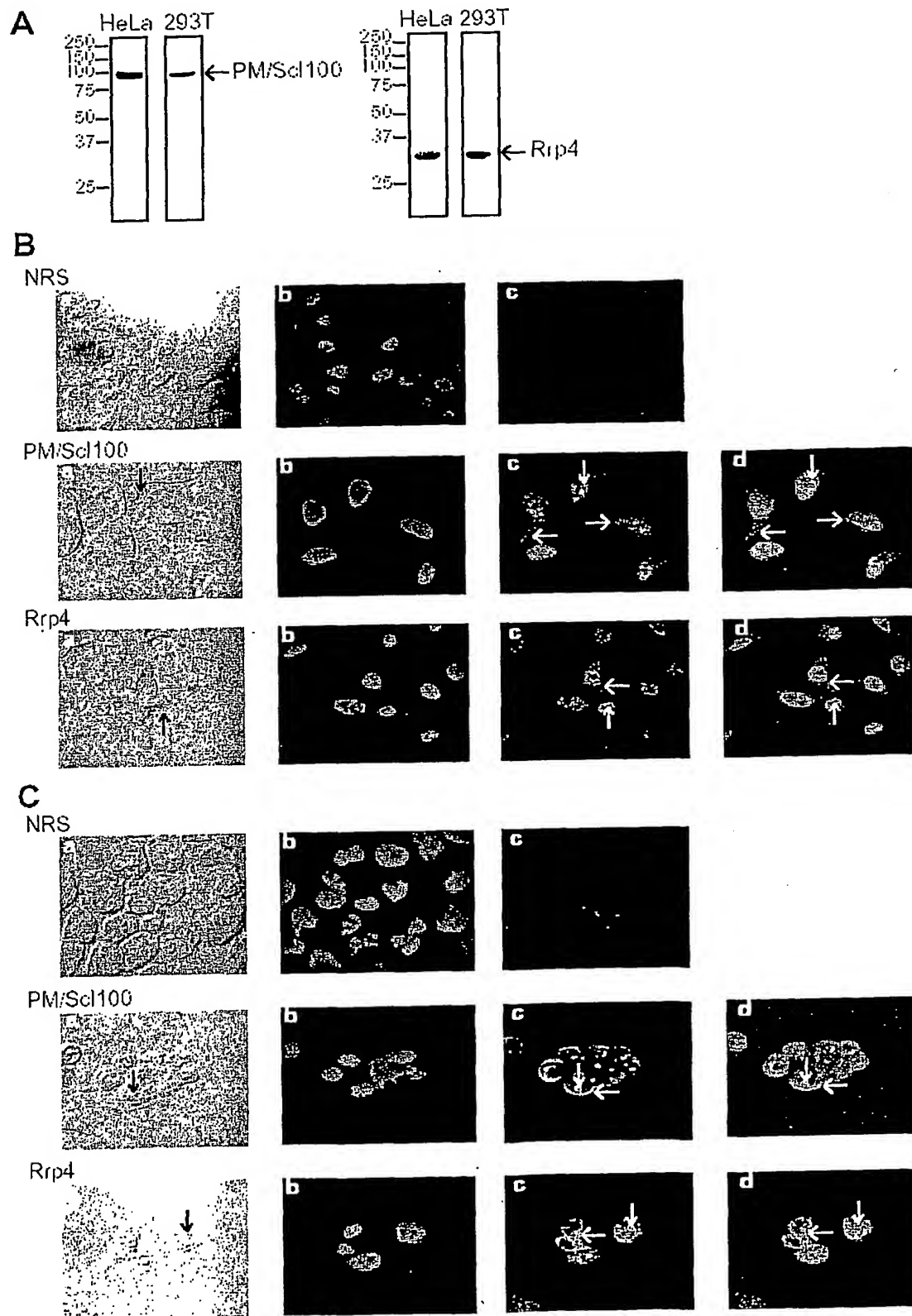


FIG. 23

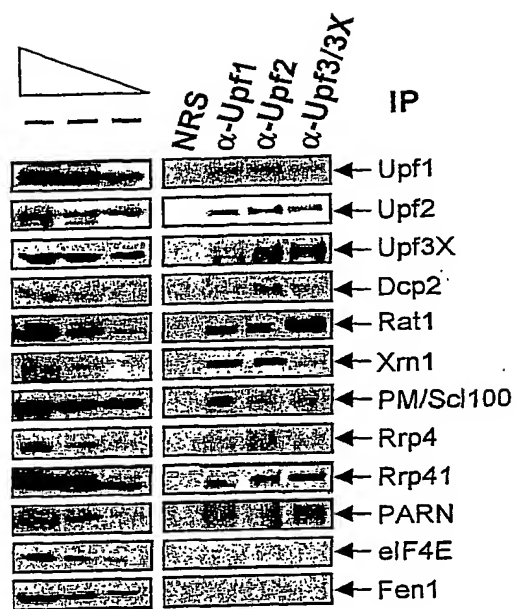


FIG. 24

FIG. 25

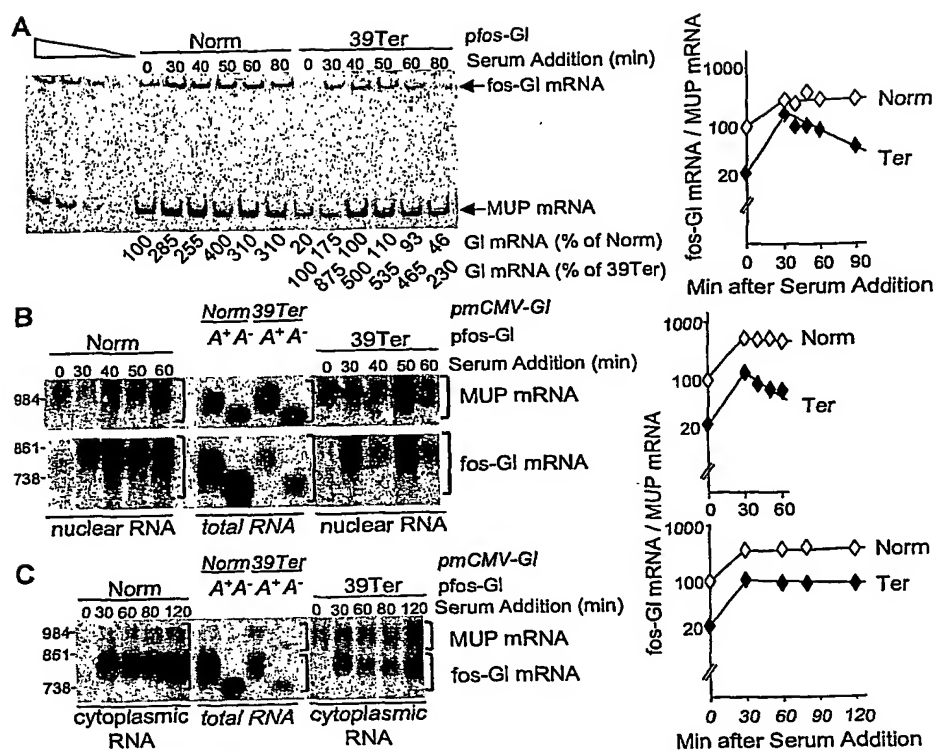


FIG. 26

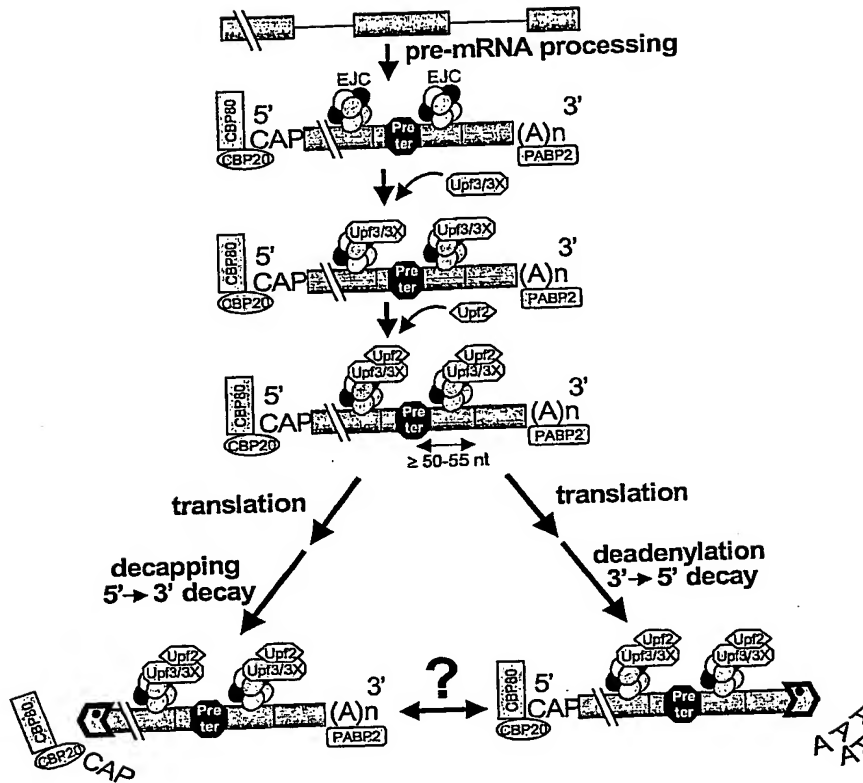
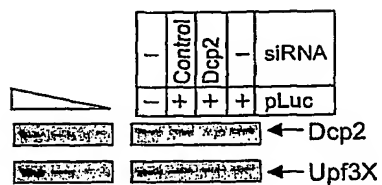
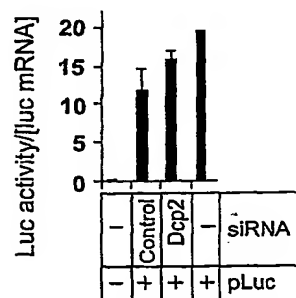
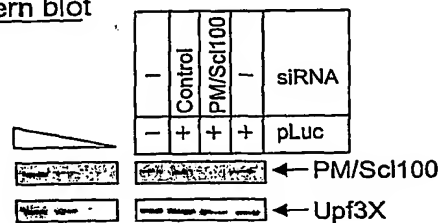
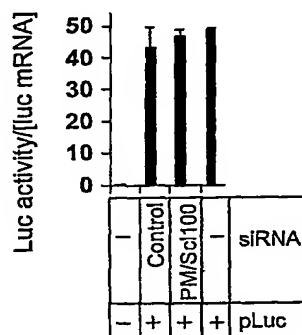
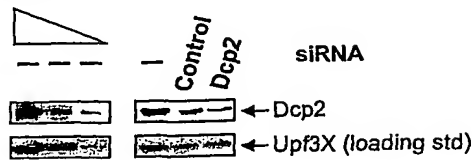
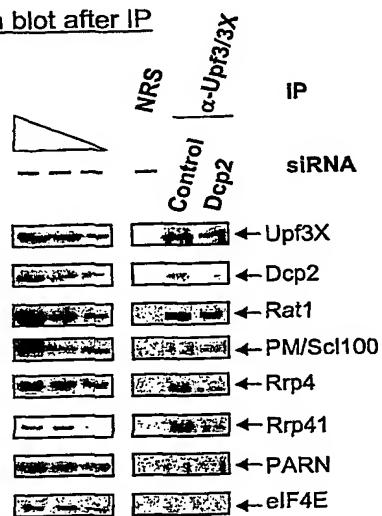
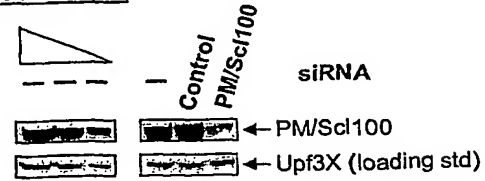
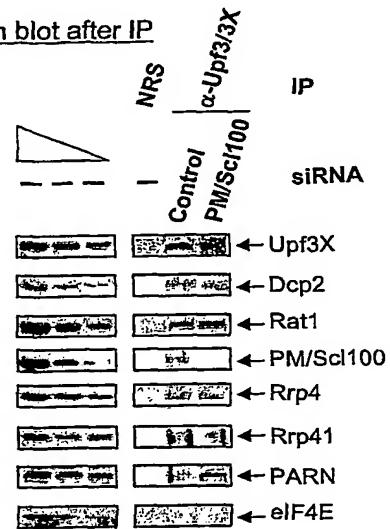
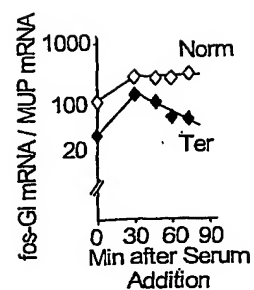
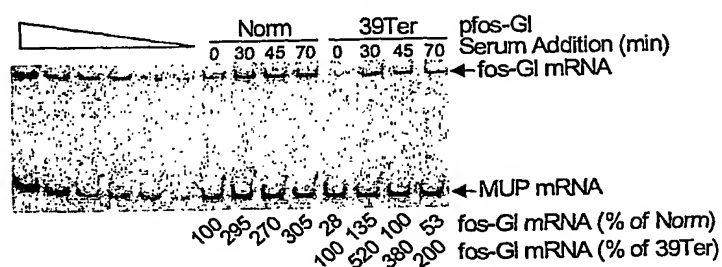
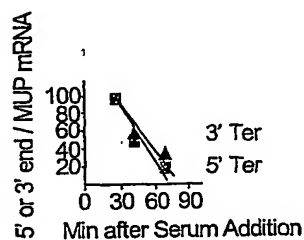


FIG. 27

A. Down-regulation of Dcp2Western blotLuciferase assay**B. Down-regulation of PM/Sci100**Western blotLuciferase assay**FIG. 28**

A. Down-regulation of Dcp2Western blotWestern blot after IP**B. Down-regulation of PM/Scf100**Western blotWestern blot after IP**FIG. 29**

A. Serum induction of pfos-Gl mRNA**B. Comparison of fos-Gl 39Ter mRNA 5' and 3' ends**

time end	30'	45'	70'
3' Ter	100	61	38
5' Ter	100	52	23

FIG. 30

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